

GenCore version 5.1.4_p5_4578
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OM nucleic - nucleic search, using sw model

Run on: March 11, 2003, 09:15:13 : Search time 323.659 Seconds
(without alignments)
70.164 Million cell updates/sec

Title: US-09-913-524-32

Perfect score: 34
Sequence: 1 aggcctcggaggacccgctgcctgcgaact 34

Scoring table:
IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 478924 seqs, 33359956 residues

Total number of hits satisfying chosen parameters: 957848

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications_NA:*

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match Length | ID | Description |
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| 1 | 20.4 | 60.0 | 247 9 | US-09-796-692-3875 |
| 2 | 20.4 | 60.0 | 1125 10 | US-09-925-300-674 |
| 3 | 20.2 | 59.4 | 1812 10 | US-09-750-240-3 |
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| 5 | 20.2 | 59.4 | 3192 9 | US-10-121-049-75 |
| 6 | 20.2 | 59.4 | 3192 9 | US-10-123-904-75 |
| 7 | 20.2 | 59.4 | 3192 9 | US-10-140-470-75 |
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| 9 | 20.2 | 59.4 | 3192 9 | US-10-176-918-75 |
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| 12 | 20.2 | 59.4 | 3192 9 | US-10-140-474-75 |
| 13 | 20.2 | 59.4 | 3192 9 | US-10-142-431-75 |
| 14 | 20.2 | 59.4 | 3192 9 | US-10-143-114-75 |
| 15 | 20.2 | 59.4 | 3192 9 | US-10-140-002-75 |
| 16 | 20.2 | 59.4 | 3549 10 | US-09-750-240-5 |
| 17 | 20.2 | 59.4 | 3552 10 | US-09-750-240-10 |
| 18 | 20.2 | 59.4 | 4942 9 | US-10-201-000-1 |
| 19 | 19.2 | 56.5 | 703 10 | US-09-822-830A-532 |

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| 20 | 19.2 | 56.5 | 4736 9 | US-09-764-868-1491 | Sequence 1491, Ap |
| 21 | 18.8 | 55.3 | 155074 9 | US-10-026-188-6 | Sequence 6, Appl1 |
| 22 | 18.6 | 54.7 | 554 9 | US-10-046-935-164 | Sequence 164, App |
| 23 | 18.6 | 54.7 | 554 9 | US-09-878-178-164 | Sequence 164, App |
| 24 | 18.6 | 54.7 | 777 10 | US-09-939-825-12 | Sequence 12, Appl |
| 25 | 18.6 | 54.7 | 1770 10 | US-09-972-086-3 | Sequence 3, Appl1 |
| 26 | 18.6 | 54.7 | 4473 10 | US-09-751-100B-1 | Sequence 1, Appl1 |
| 27 | 18.6 | 54.7 | 4593 10 | US-09-972-086-1 | Sequence 1, Appl1 |
| 28 | 18.4 | 54.1 | 176 10 | US-09-876-889-331 | Sequence 331, Appl |
| 29 | 18.4 | 54.1 | 555 10 | US-09-764-860-235 | Sequence 235, App |
| 30 | 18.4 | 54.1 | 5775 9 | US-09-866-557A-1 | Sequence 1, Appl1 |
| 31 | 18.4 | 54.1 | 14654 10 | US-09-764-860-1054 | Sequence 1054, Ap |
| 32 | 18.2 | 53.5 | 1308 10 | US-09-801-368-149 | Sequence 149, App |
| 33 | 18.2 | 53.5 | 1338 9 | US-09-796-753-129 | Sequence 129, App |
| 34 | 18.2 | 53.5 | 1347 9 | US-09-974-879-39 | Sequence 39, Appl |
| 35 | 18.2 | 53.5 | 1368 10 | US-09-815-242-7684 | Sequence 7684, Ap |
| 36 | 18.2 | 53.5 | 2756 10 | US-09-764-853-898 | Sequence 898, App |
| 37 | 18.2 | 53.5 | 2940 9 | US-10-171-229-1 | Sequence 1, Appl1 |
| 38 | 18.2 | 53.5 | 15425 10 | US-09-764-869-1654 | Sequence 1654, Ap |
| 39 | 18.2 | 53.5 | 15613 10 | US-09-764-877-3525 | Sequence 3525, App |
| 40 | 18 | 52.9 | 470 10 | US-09-960-352-2406 | Sequence 2406, Ap |
| 41 | 18 | 52.9 | 970 9 | US-09-822-846-546 | Sequence 546, App |
| 42 | 18 | 52.9 | 1013 10 | US-09-822-849A-226 | Sequence 226, App |
| 43 | 18 | 52.9 | 1083 10 | US-09-974-300-1692 | Sequence 1692, Ap |
| 44 | 18 | 52.9 | 2753 9 | US-09-774-639-12 | Sequence 12, Appl |
| 45 | 18 | 52.9 | 4701 10 | US-09-864-864-326 | Sequence 326, App |

ALIGNMENTS

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RESULT 1
US-09-796-692-3875/c
: Sequence 3875, Application US/09796692
: Publication No. US20020198362A1
GENERAL INFORMATION:
: APPLICANT: Galger, Alexander
: APPLICANT: Mamillon, Jane
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DETECTION, DIAGNOSIS AND TH
: TITLE OF INVENTION: HEMATOLOGICAL MALIGNANCIES
: FILE REFERENCE: 2077.001200
: CURRENT APPLICATION NUMBER: US/09/79 692
: CURRENT FILING DATE: 2001-03-01
: PRIOR APPLICATION NUMBER: 60/186,126
: PRIOR FILING DATE: 2000-03-01
: PRIOR APPLICATION NUMBER: 60/190,479
: PRIOR FILING DATE: 2000-03-17
: PRIOR APPLICATION NUMBER: 60/200,545
: PRIOR FILING DATE: 2000-04-27
: PRIOR APPLICATION NUMBER: 60/200,303
: PRIOR FILING DATE: 2000-04-28
: PRIOR APPLICATION NUMBER: 60/200,779
: PRIOR FILING DATE: 2000-04-28
: PRIOR APPLICATION NUMBER: 60/200,999
: PRIOR FILING DATE: 2000-05-01
: PRIOR APPLICATION NUMBER: 60/202,084
: PRIOR FILING DATE: 2000-05-04
: PRIOR APPLICATION NUMBER: 60/206,201
: PRIOR FILING DATE: 2000-05-22
: PRIOR APPLICATION NUMBER: 60/218,950
: PRIOR FILING DATE: 2000-07-14
: PRIOR APPLICATION NUMBER: 60/222,903
: PRIOR FILING DATE: 2000-08-03
: PRIOR APPLICATION NUMBER: 60/223,416
: PRIOR FILING DATE: 2000-08-04
: PRIOR APPLICATION NUMBER: 60/223,378
: PRIOR FILING DATE: 2000-08-07
: NUMBER OF SEQ ID NOS: 9597
: SOFTWARE: FastSeq for Windows Version 3.0
: SEQ ID NO 3875
: LENGTH: 247
: TYPE: DNA
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: ORGANISM: Homo sapiens
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Best Local Similarity 80.0%; Score 20.4; DB 9; Length 247;
Matches 24; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

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RESULT 2
US-09-925-300-674
: Sequence 674, Application US/09925300
: Patent No. US20020151681A1
: GENERAL INFORMATION:
: APPLICANT: Craig Rosen,
: APPLICANT: Steve Ruben,
: TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
: FILE REFERENCE: PA101
: CURRENT APPLICATION NUMBER: US/09/925,300
: CURRENT FILING DATE: 2001-08-10
: PRIOR APPLICATION NUMBER: PCT/US00/05988
: PRIOR FILING DATE: 2000-03-08
: PRIOR APPLICATION NUMBER: 60/124,270
: PRIOR FILING DATE: 1999-03-12
: NUMBER OF SEQ ID NOS: 1890
: SOFTWARE: PatentIn Ver. 2.0
: SEQ ID NO 674
: LENGTH: 1125
: TYPE: DNA
: ORGANISM: Homo sapiens
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US-09-925-300-674

Query Match
Best Local Similarity 80.0%; Score 20.4; DB 10; Length 1125;
Matches 24; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

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RESULT 3
US-09-750-240-3
: Sequence 3, Application US/09750240
: Patent No. US20020103147A1
: GENERAL INFORMATION:
: APPLICANT: Hammon, H. K.,
: APPLICANT: Jinsel, P. A.,
: APPLICANT: Ping, P.,
: APPLICANT: Post, S. R.,
: APPLICANT: Gao, M.,
: TITLE OF INVENTION: GENE THERAPY FOR CONGESTIVE HEART
: FILE REFERENCE: 220002056723
: CURRENT APPLICATION NUMBER: US/09/750,240
: CURRENT FILING DATE: 2001-10-12
: PRIOR APPLICATION NUMBER: US 09/472,667
: PRIOR FILING DATE: 1999-12-27
: PRIOR APPLICATION NUMBER: US 09/008,097
: PRIOR FILING DATE: 1998-01-16
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: PRIOR APPLICATION NUMBER: US 08/924,757
: PRIOR FILING DATE: 1997-09-05
: PRIOR APPLICATION NUMBER: US 60/048,933
: PRIOR FILING DATE: 1997-06-16
: PRIOR APPLICATION NUMBER: US 08/708,600
: PRIOR FILING DATE: 1996-09-05
: NUMBER OF SEQ ID NOS: 13
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 3
: LENGTH: 1812
: TYPE: DNA
: ORGANISM: Homo sapiens
US-09-750-240-3

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Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

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Db 668 GGCTCCGAGACCGGCTGCCATGCCACT 700

RESULT 4
US-10-028-072-75
: Sequence 75, Application US/10028072
: Publication No. US20030004311A1
: GENERAL INFORMATION:
: APPLICANT: Baker, Kevin P.,
: APPLICANT: Beresini, Maureen
: APPLICANT: Deforge, Laura
: APPLICANT: Desnoyers, Luc
: APPLICANT: Flivaroff, Ellen
: APPLICANT: Gao, Wei-Qiang
: APPLICANT: Gerritsen, Mary E.,
: APPLICANT: Goddard, Audrey
: APPLICANT: Godowski, Paul J.,
: APPLICANT: Gurney, Austin L.,
: APPLICANT: Sherwood, Steven
: APPLICANT: Smith, Victoria
: APPLICANT: Stewart, Timothy A.,
: APPLICANT: Tumas, Daniel
: APPLICANT: Watanabe, Colin K
: APPLICANT: Wood, William
: APPLICANT: Zhang
: TITLE OF INVENTION:
: FILE REFERENCE:
: CURRENT APPLICATION NUMBER: US/10/028,072
: CURRENT FILING DATE: 2001-12-19
: PRIOR APPLICATION NUMBER: 60/049911
: PRIOR FILING DATE: 1997-06-18
: PRIOR APPLICATION NUMBER: 60/056974
: PRIOR FILING DATE: 1997-08-26
: PRIOR APPLICATION NUMBER: 60/059113
: PRIOR FILING DATE: 1997-09-17
: PRIOR APPLICATION NUMBER: 60/059115
: PRIOR FILING DATE: 1997-09-17
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: PRIOR FILING DATE: 1997-09-17
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: PRIOR FILING DATE: 1997-09-17
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: PRIOR FILING DATE: 1997-09-17
: PRIOR APPLICATION NUMBER: 60/059263
: PRIOR FILING DATE: 1997-09-18
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: PRIOR FILING DATE: 1997-09-19
: PRIOR APPLICATION NUMBER: 60/059588
: PRIOR FILING DATE: 1997-09-19
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: PRIOR FILING DATE: 1997-09-24
: PRIOR APPLICATION NUMBER: 60/062250
: PRIOR FILING DATE: 1997-10-17
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; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
```

```
Query Match          59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
```

```
OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAC 33
DB 11 ATGGCTCGSAGAGCTCGGTTGCCCATCCGAGC 43
```

RESULT 5

```
US-10-121-049-75
; Sequence 75, Application US/10121049
; Publication No. US20030022339A1
```

GENERAL INFORMATION:

```
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C17
; CURRENT APPLICATION NUMBER: US/10/121,049
; CURRENT FILING DATE: 2002-04-12
; Prior Application removed - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 75
; LENGTH: 3192
; TYPE: DNA
; ORGANISM: Homo Sapien
; FEATURE:
; NAME/KEY: unsure
; LOCATION: 1428, 1431
; OTHER INFORMATION: unknown base
US-10-121-049-75
```

```
Query Match          59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
```

```
OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAC 33
DB 11 ATGGCTCGSAGAGCTCGGTTGCCCATCCGAGC 43
```

RESULT 6

```
US-10-123-904-75
; Sequence 75, Application US/10123904
; Publication No. US20030022328A1
```

GENERAL INFORMATION:

```
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
```

```
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C54
; CURRENT APPLICATION NUMBER: US/10/123,904
; CURRENT FILING DATE: 2002-04-16
; Prior Application removed - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 75
; LENGTH: 3192
; TYPE: DNA
; ORGANISM: Homo Sapien
; FEATURE:
; NAME/KEY: unsure
; LOCATION: 1428, 1431
; OTHER INFORMATION: unknown base
US-10-123-904-75
```

```
Query Match          59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
```

```
OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAC 33
DB 11 ATGGCTCGSAGAGCTCGGTTGCCCATCCGAGC 43
```

RESULT 7

```
US-10-140-470-75
; Sequence 75, Application US/10140470
; Publication No. US20030022331A1
```

GENERAL INFORMATION:

```
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C160
; CURRENT APPLICATION NUMBER: US/10/140,470
; CURRENT FILING DATE: 2002-05-06
; Prior Application removed - See Palm or File Wrapper
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 75
; LENGTH: 3192
; TYPE: DNA
; ORGANISM: Homo Sapien
; FEATURE:
```

NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-140-470-75

Query Match
Best Local Similarity 59.4%; Score 20.2; DB 9; Length 3192;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAG 33
DB 11 ATGGCTCCGAGAGCTCCGCTTGCCCATGCCAGC 43

RESULT 8
US-10-175-746-75

Sequence 75, Application US/10175746
Publication No. US20030027270A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Elivaroff, Ellen
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C353
CURRENT APPLICATION NUMBER: US/10/175,746
CURRENT FILING DATE: 2002-06-15
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-175-746-75

Query Match
Best Local Similarity 59.4%; Score 20.2; DB 9; Length 3192;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAG 33
DB 11 ATGGCTCCGAGAGCTCCGCTTGCCCATGCCAGC 43

RESULT 9
US-10-176-918-75

Sequence 75, Application US/10176918
Publication No. US20030027275A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Elivaroff, Ellen
APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C382
CURRENT APPLICATION NUMBER: US/10/176,918
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-176-918-75

Query Match
Best Local Similarity 59.4%; Score 20.2; DB 9; Length 3192;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGAGACCGCTGCCCATGCCAAG 33
DB 11 ATGGCTCCGAGAGCTCCGCTTGCCCATGCCAGC 43

RESULT 10
US-10-176-921-75

Sequence 75, Application US/10176921
Publication No. US20030027276A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Elivaroff, Ellen
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C288
CURRENT APPLICATION NUMBER: US/10/176,921
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431

OTHER INFORMATION: unknown base
US-10-176-921-75

Query Match 59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGACCGCTGCCATGCGAC 33
DB 11 ATGGCTCCGAGGCTCCGCTTGCCATCGAGC 43

RESULT 11

US-10-137-865-75

Sequence 75, Application US/10137865
Publication No. US20030032155A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Filvaroff, Ellen
APPLICANT: Gao, Mei-Qiang
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C154
CURRENT APPLICATION NUMBER: US/10/137,865
CURRENT FILING DATE: 2002-05-03
Prior Application removed - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-137-865-75

Query Match 59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGACCGCTGCCATGCGAC 33
DB 11 ATGGCTCCGAGGCTCCGCTTGCCATCGAGC 43

RESULT 12

US-10-140-474-75

Sequence 75, Application US/10140474
Publication No. US20030032156A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Filvaroff, Ellen
APPLICANT: Gao, Mei-Qiang
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey

APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C162
CURRENT APPLICATION NUMBER: US/10/140,474
CURRENT FILING DATE: 2002-05-06
Prior Application removed - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-140-474-75

Query Match 59.4%; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.8%; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

OY 1 AGGCTCCGAGACCGCTGCCATGCGAC 33
DB 11 ATGGCTCCGAGGCTCCGCTTGCTGATCGAGC 43

RESULT 13

US-10-142-431-75

Sequence 75, Application US/10142431
Publication No. US20030036179A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Deforge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Filvaroff, Ellen
APPLICANT: Gao, Mei-Qiang
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3330R1C251
CURRENT APPLICATION NUMBER: US/10/142,431
CURRENT FILING DATE: 2002-05-10
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 75
LENGTH: 3192
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 1428, 1431
OTHER INFORMATION: unknown base
US-10-142-431-75

Query Match 59.48; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.88; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY 1 AGGCTCCGAGAGACCGGCTGCCATGCCAC 33
DB 11 ATGGCTCCGAGAGCTCCGTTGCCATCCGAGC 43

RESULT 14

US-10-143-114-75

; Sequence 75, Application US/10143114
; Publication No. US20030036180A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3330R1C211
; CURRENT APPLICATION NUMBER: US/10/143,114
; CURRENT FILING DATE: 2002-05-09
; Prior Application removed - See Palm or File Wrapper
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 75
; LENGTH: 3192
; TYPE: DNA
; ORGANISM: Homo Sapien
; FEATURE:
; NAME/KEY: unsure
; LOCATION: 1428, 1431
; OTHER INFORMATION: unknown base
US-10-143-114-75

Query Match 59.48; Score 20.2; DB 9; Length 3192;

Best Local Similarity 75.88; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY 1 AGGCTCCGAGAGACCGGCTGCCATGCCAC 33
DB 11 ATGGCTCCGAGAGCTCCGTTGCCATCCGAGC 43

RESULT 15

US-10-140-002-75

; Sequence 75, Application US/10140002
; Publication No. US20030037623A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.

; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3330R1C59
; CURRENT APPLICATION NUMBER: US/10/140,002
; CURRENT FILING DATE: 2002-05-06
; Prior Application removed - See Palm or File Wrapper
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 75
; LENGTH: 3192
; TYPE: DNA
; ORGANISM: Homo Sapien
; FEATURE:
; NAME/KEY: unsure
; LOCATION: 1428, 1431
; OTHER INFORMATION: unknown base
US-10-140-002-75

Query Match 59.48; Score 20.2; DB 9; Length 3192;
Best Local Similarity 75.88; Pred. No. 52;
Matches 25; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY 1 AGGCTCCGAGAGACCGGCTGCCATGCCAC 33
DB 11 ATGGCTCCGAGAGCTCCGTTGCCATCCGAGC 43

Search completed: March 11, 2003, 14:20:01
Job time : 327.659 secs